

Master in Artificial Intelligence



Algorithm Selection & Development V





Purpose

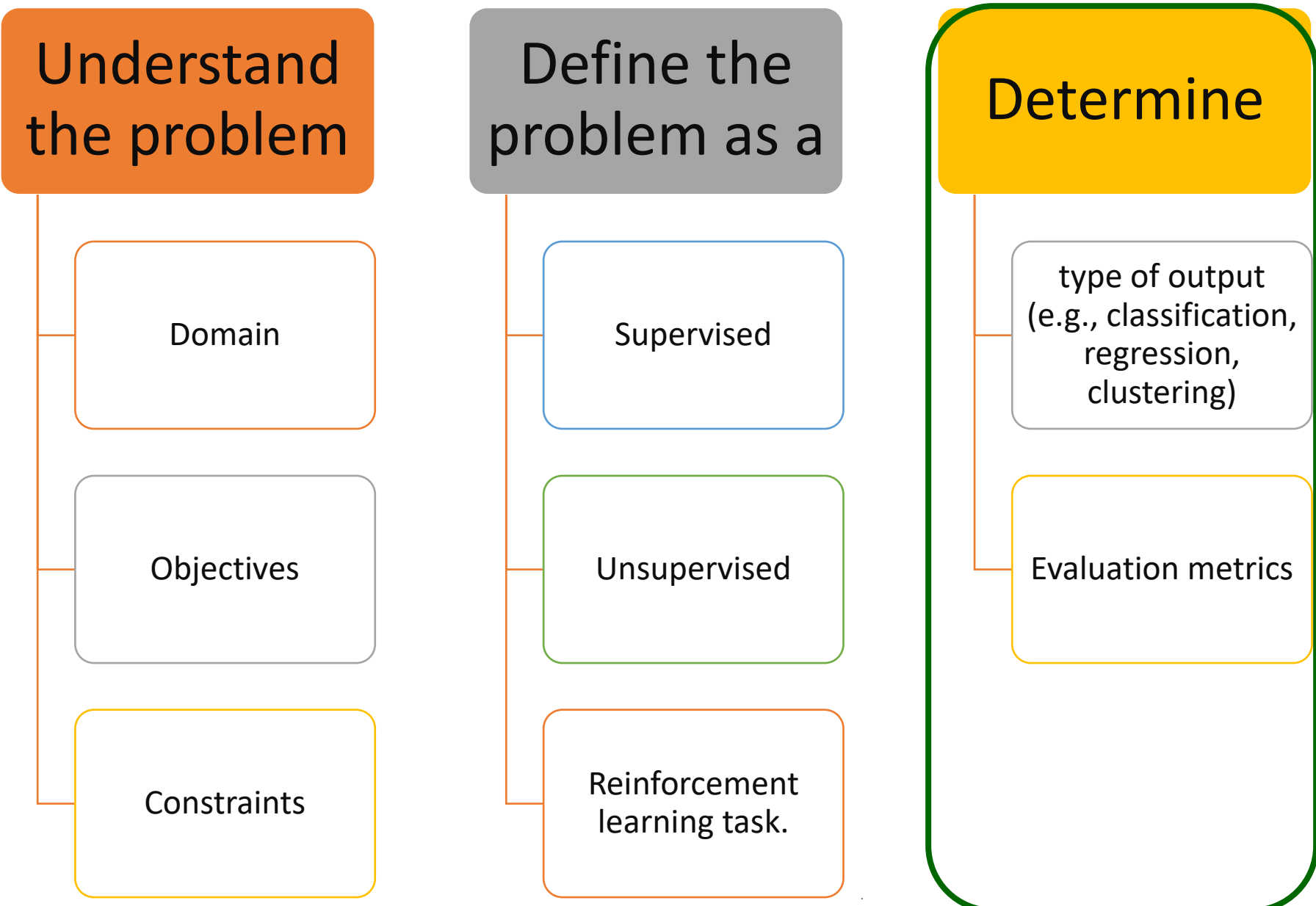
The purpose of the section is to help you learn how to research, select, and develop appropriate algorithms to become a Successful Artificial Intelligence (AI) Engineer

At the end of this lecture, you will learn the following

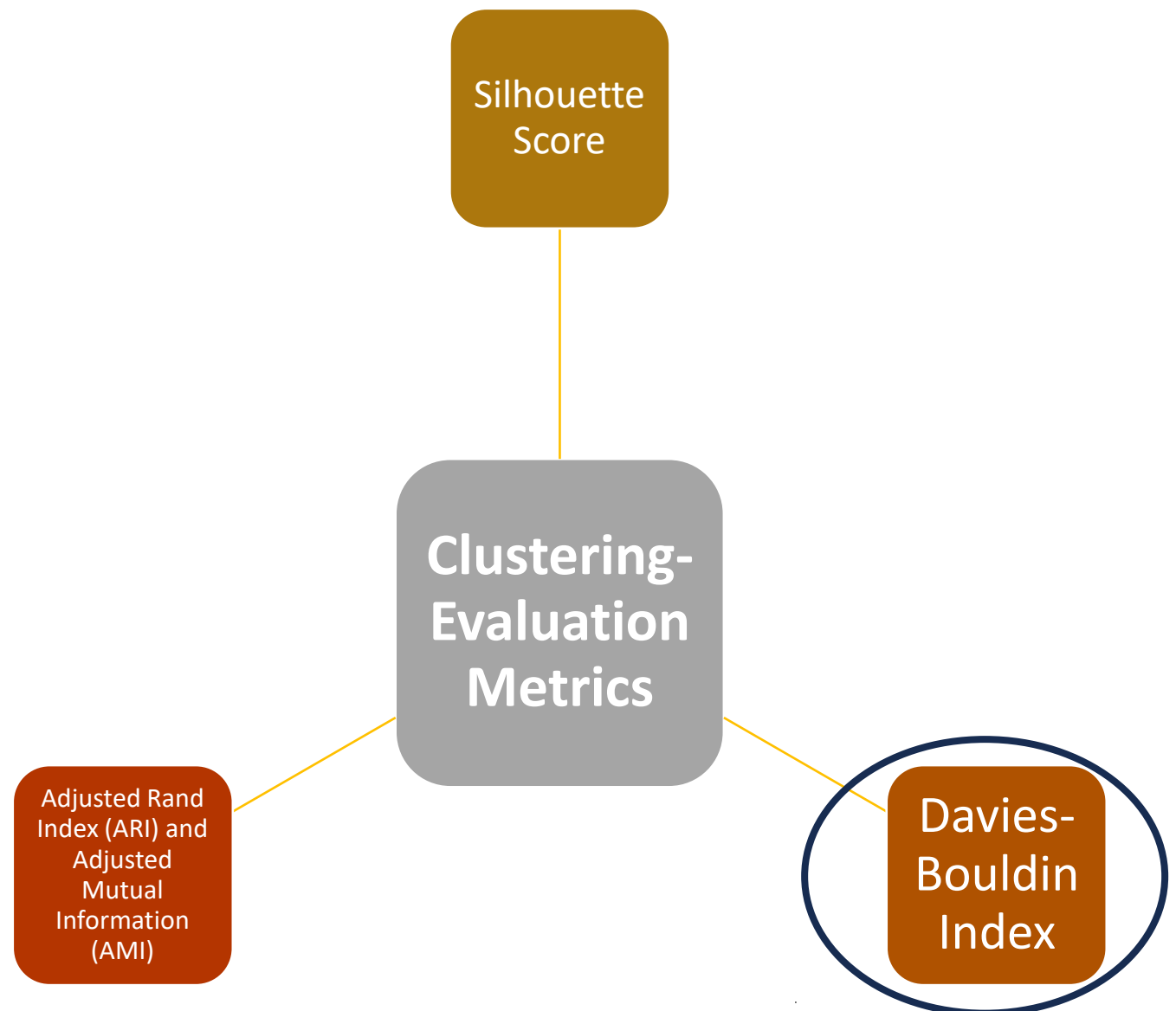
- **How does Davies-Bouldin Index compute the average similarity between each cluster and its most similar cluster**



How to determine type of output and evaluation metrics?

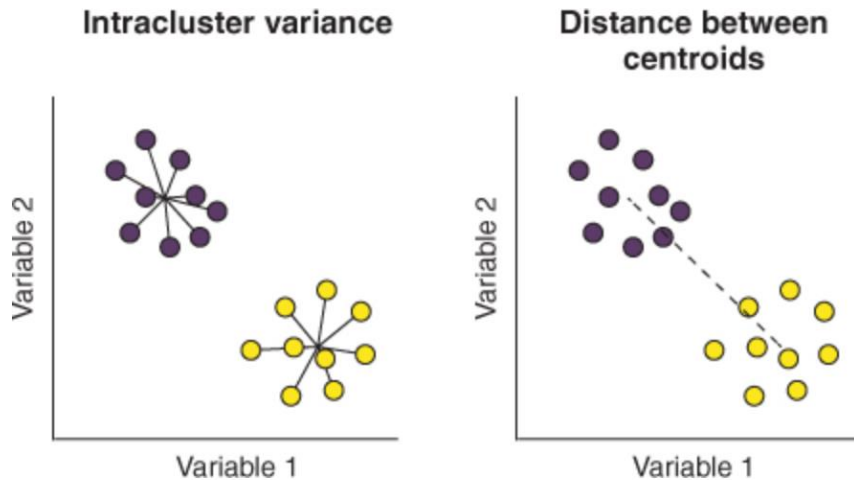


How does Davies-Bouldin Index compute the average similarity between each cluster and its most similar cluster



Calculating the Davies-Bouldin Index

- For each cluster i :
 - Compute the centroid C_i , which represents the center of the cluster.
 - For each other cluster j (where $j \neq i$):
 - Calculate the average distance between each point in cluster i and the centroid C_i . This is denoted as avg_i .
 - Calculate the average distance between each point in cluster j and the centroid C_j . This is denoted as avg_j .
 - Compute the distance between the centroids C_i and C_j . This is denoted as d_{ij} .
 - Calculate the similarity between cluster i and its most similar cluster j as
$$R_{ij} = (avg_i + avg_j) / d_{ij}$$
- The Davies-Bouldin Index for the clustering is the average of the similarity values R_{ij} across all clusters



Interpreting the Davies-Bouldin Index

Low Davies-Bouldin Index

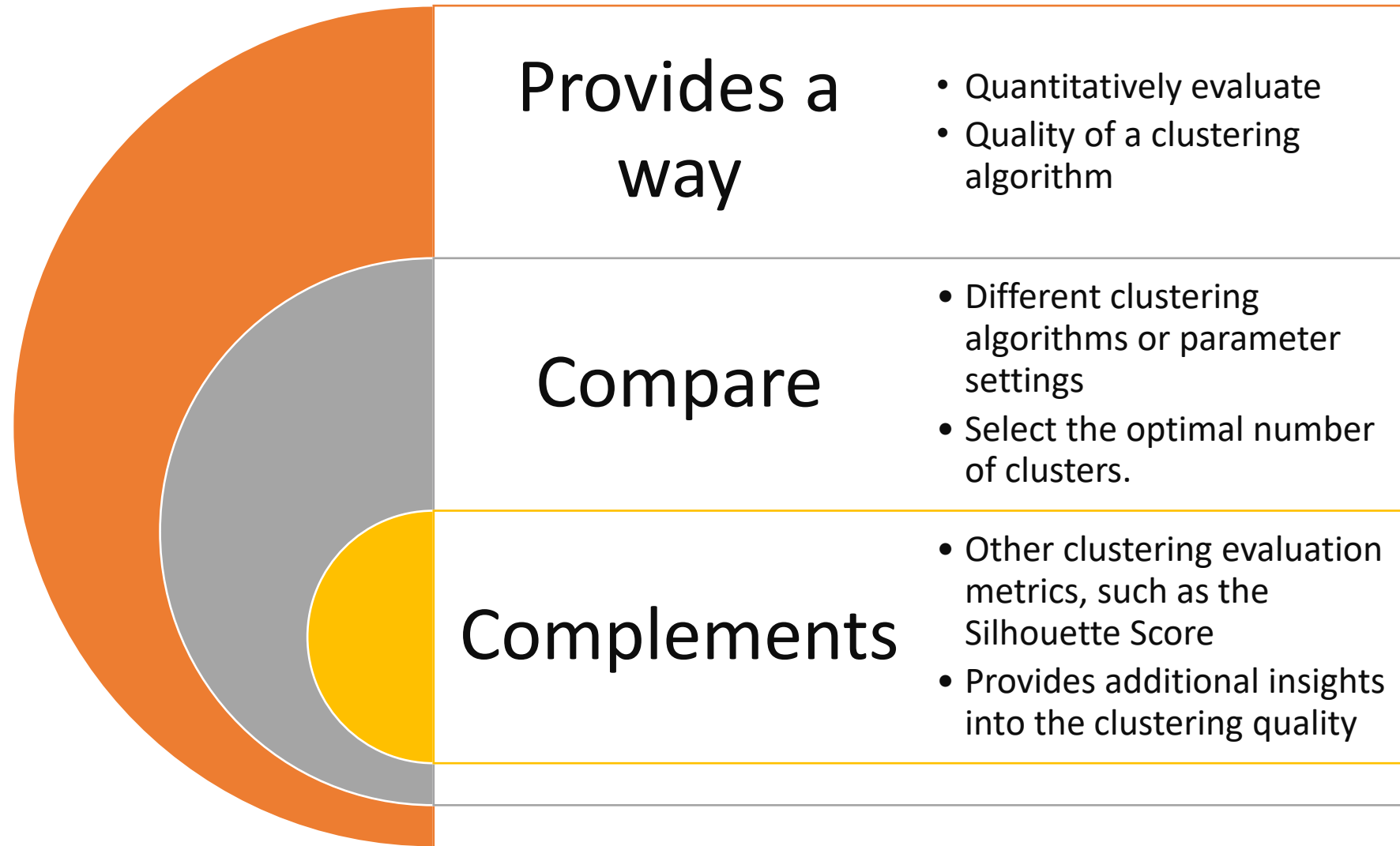
- Clusters are well-separated and distinct from each other
- Each cluster is more similar to its own cluster than to others

High Davies-Bouldin Index

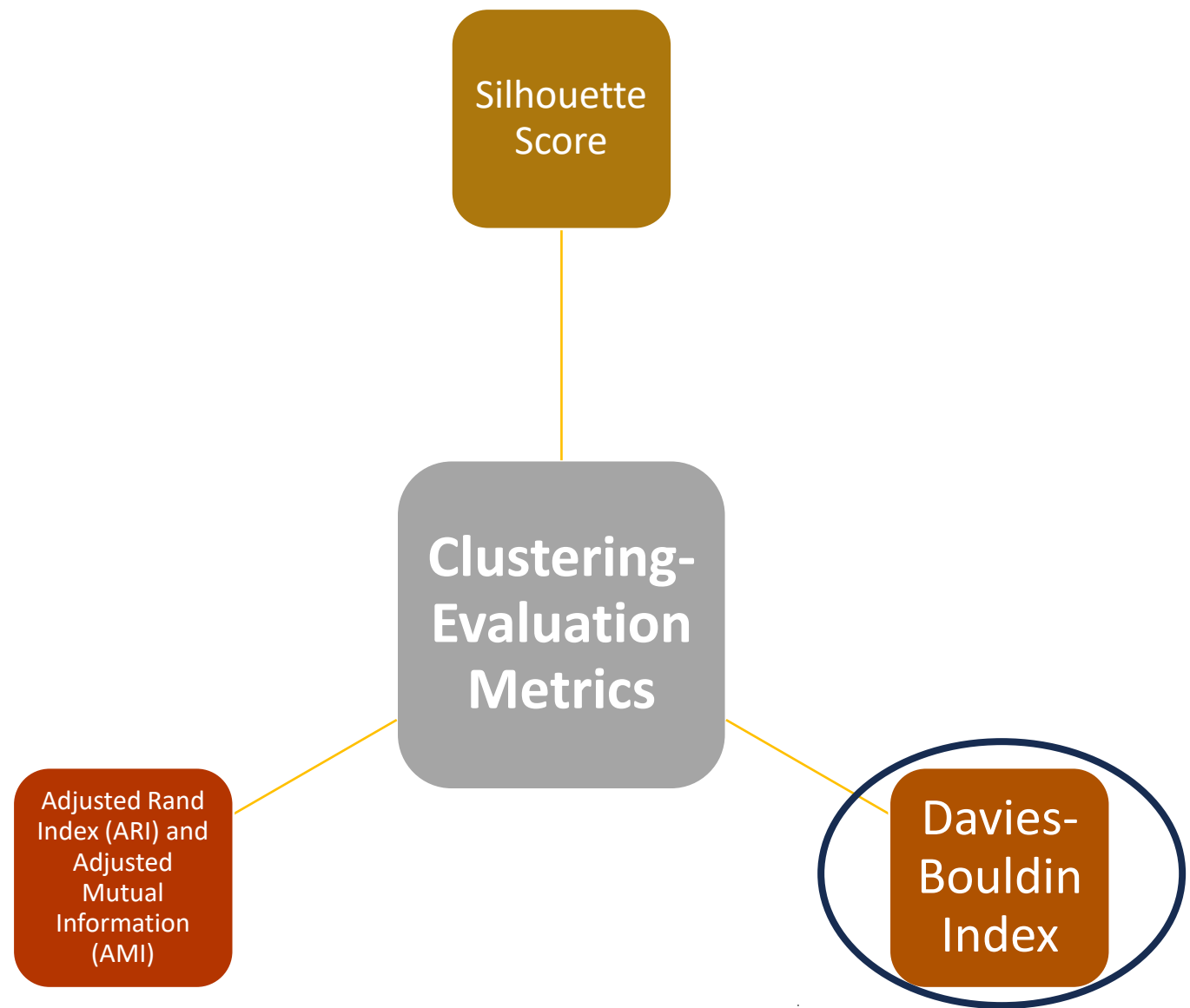
- Clusters are less well-separated and may have significant overlap
- The boundaries between clusters are less clear



Using the Davies-Bouldin Index for Cluster Evaluation

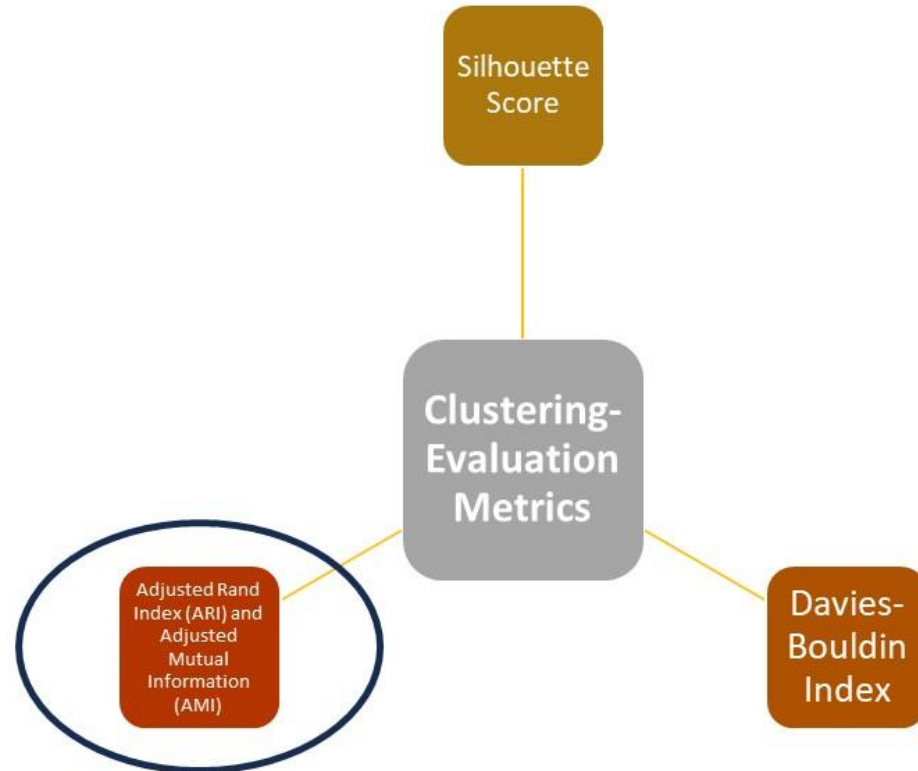


How does Davies-Bouldin Index compute the average similarity between each cluster and its most similar cluster



What is next?

Adjusted Rand Index (ARI) and Adjusted Mutual Information (AMI) Measures the agreement between true labels and cluster assignments



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